

22-25 November 2015

Chatrium Hotel Riverside, Bangkok, Thailand

"The Next-Generation Computational Biology"

With the advancement in next-generation molecular technology in generating the high-throughput "omics" data, life science has come to the era of "Big Data". To extract biological knowledge from the data, and translate it into benefits for society (e.g. better medicine and healthcare), novel and advanced computational tools are needed for data analysis. CSBio 2015 would like to invite researchers and industrial counterparts to meet at this event to exchange ideas and stimulate research collaborations.

CALL FOR PAPERS

We invite you to submit either abstracts, extended abstracts, and full papers as the following choices:

1. Poster presentation: Abstracts
2. Conference proceedings: Extended abstracts (not exceed 5 pages)
3. Full research papers to be considered for
 - Journal of Bioinformatics and Computational Biology (JBCB)
 - Computational and Structural Biotechnology Journal (CSBJ)

TOPICS include (but are not limited to)

Adaptive computation in bioinformatics
Bio-data visualization
Bio-inspired computing
Biological network reconstruction and analysis
Biomarker discovery
Disease classification
DNA, RNA and protein sequence analysis
Drug discovery and validation
Epigenetics or epigenomics
Functional genomics

Gene expression analysis
Medical and biomedical informatics
Modeling and simulation of biological processes, pathways, etc.
Molecular evolution and phylogeny
Next-generation sequencing
Parallel and distributed computing for life science
Population genetics
Proteomics and other omics
Protein folding
Translational bioinformatics

IMPORTANT DATES

Paper submission deadline	31 July 2015 (Extended)
Notification date	15 September 2015
Early-bird registration deadline	22 September 2015

KEYNOTE SPEAKERS

Steve Rozen, Ph.D.

Duke-NUS Graduate medical school, Singapore

On *"Genome bioinformatics' current successes, challenges, and opportunities in the era of low-cost sequencing"*

Saman K. Halgamuge, Ph.D.

University of Melbourne, Australia

On *"Big data analysis in Metagenomics"*

WORKSHOPS

"Next-Generation Sequencing Data Analysis" by

Weerayuth Kittichotirat, Ph.D. and

Teeraphan Laomettachit, Ph.D.

KMUTT, Thailand

"Algorithms for Detecting Protein Complexes from Whole-Genome PPI Networks" by

Leong Hon Wai, Ph.D.

National University of Singapore, Singapore

INVITED SPEAKERS

Panagiotou Gianni, Ph.D.

The University of Hong Kong, Hong Kong

On *"Microbiome systems biology"*

Sissades Tongsim, Ph.D.

BIOTEC, Thailand

On *"Complex population substructure and genome wide association studies"*

Philip Shaw, Ph.D.

BIOTEC, Thailand

On *"RNA-seq methods for studying the malarial transcriptome"*

Prapat Suriyaphol, Ph.D.

Mahidol University, Thailand

On *"Precision medicine: The future of healthcare"*

For more information:



<http://www.csbio.org/>